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=> HCV
         14830 HCV
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L1
                 (HCV OR HCVS)
=> mutation (1) 2204
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           432 2204
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L2
=> mutation (L) 1067
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L3
            21 MUTATION (L) 1067
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=> L2 and L4
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L10
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=> L2 and L5

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=> L2 and L6

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=> L2 and L7

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=> L1 and L3

L14 1 L1 AND L3

=> L1 and L4

2 L1 AND L4 L15

=> L1 and L5

1 L1 AND L5 L16

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The search profile that was entered contains terms or nested terms that are not separated by a logical operator.

=> L1 and L6

L17 1 L1 AND L6

=> D L15 IBIB ABS 1-2

L15 ANSWER 1 OF 2 CAPLUS COPYRIGHT 2008 ACS on STN

2005:523226 CAPLUS ACCESSION NUMBER:

DOCUMENT NUMBER: 143:54458

TITLE: Replication competent hepatitis C virus genotype 1a

with adaptive mutations and methods of use for drug

screening and selection of host cell line

INVENTOR(S): Lemon, Stanley M.; Yi, Minkyung

Board of Regents, the University of Texas System, USA PATENT ASSIGNEE(S):

SOURCE: PCT Int. Appl., 102 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent English LANGUAGE:

FAMILY ACC. NUM. COUNT: 1

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PRIORITY APPLN. INFO.:
                                                 US 2003-525989P
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                                                 WO 2004-US40120
                                                                       W 20041201
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AB The invention provides replication competent polynucleotides that include a coding sequence encoding a hepatitis C virus polyprotein having adaptive mutations. The genotype 1a adaptive mutations identified here can be grouped functionally into two groups: K2040R, F2080V, and S2204I, which are all located within NS5A, and Q1067R, G1188R, V1655I, and K1691R (in NS4A), which are all located in or associated with the protease domain of NS3. These NS3 and NS4A mutations are located at some distance from other genotype 1a adaptive mutations in NS3 that were previously described. The contribution of the NS5A adaptive mutations to the replication of genotype 1a RNA appears to be additive to that of the NS3/4A mutations and not synergistic as shown for the combination of Q1067R and K1691R. The invention also includes methods for making replication competent polynucleotides, identifying a compound that inhibits replication of a replication competent polynucleotide, and detecting a replication competent polynucleotide, and detecting a replication competent polynucleotide.

L15 ANSWER 2 OF 2 CAPLUS COPYRIGHT 2008 ACS on STN

ACCESSION NUMBER: 2002:504935 CAPLUS

DOCUMENT NUMBER: 137:74392

TITLE: Self-replicating RNA molecule from hepatitis C virus

having adaptive mutations, and its uses in screening

assay for HCV replication inhibitors

INVENTOR(S): Kukolj, George; Pause, Arnim

PATENT ASSIGNEE(S): Boehringer Ingelheim (Canada) Ltd., Can.

SOURCE: PCT Int. Appl., 140 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

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AB The present invention relates generally to a hepatitis C virus (HCV) RNA mol. that self-replicates in appropriate cell lines, particularly to a self-replicating HCV RNA construct having an enhanced efficiency of establishing cell culture replication. A unique HCV RNA mol. is provided having an enhanced efficiency of establishing cell culture replication. Novel adaptive mutations have been identified within the HCV non-structural region that improves the efficiency of establishing persistently replicating HCV RNA in cell culture. This self-replicating polynucleotide mol. contains, contrary to all previous reports, a 5'-NTR that can be either an A as an alternative to the G already disclosed and therefore provides an alternative to existing systems comprising a self-replicating HCV RNA mol. The G-->A mutation gives rise to HCV RNA mols. that, in conjunction with mutations in the HCV non-structural region, such as the G(2042)C/Rmutations, possess greater efficiency of transduction and/or replication. The HCV RNA encoding polyprotein comprising one or more amino acid substitution selected from the group consisting of: R(1135)K; S(1148)G; S(1560)G; K(1691)R; L(1701)F; I(1984)V;T(1993)A; G(2042)C; G(2042)R; S(2404)P; L(2155)P; P(2166)L; M(2992)T; andE(1202)G is claimed. These RNA mols. when transfected in a cell line are useful for evaluating potential inhibitors of HCV replication.

=> D L9 IBIB ABS

ANSWER 1 OF 1 CAPLUS COPYRIGHT 2008 ACS on STN

ACCESSION NUMBER: 2005:523226 CAPLUS

DOCUMENT NUMBER: 143:54458

TITLE: Replication competent hepatitis C virus genotype 1a with adaptive mutations and methods of use for drug

screening and selection of host cell line

Lemon, Stanley M.; Yi, Minkyung INVENTOR(S):

Board of Regents, the University of Texas System, USA PATENT ASSIGNEE(S):

SOURCE: PCT Int. Appl., 102 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

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PRIORITY APPLN. INFO.:
                                                                W 20041201
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The invention provides replication competent polynucleotides that include AB a coding sequence encoding a hepatitis C virus polyprotein having adaptive mutations. The genotype la adaptive mutations identified here can be grouped functionally into two groups: K2040R, F2080V, and S2204I, which are all located within NS5A, and Q1067R, G1188R, V1655I, and K1691R (in NS4A), which are all located in or associated with the protease domain of NS3. These NS3 and NS4A mutations are located at some distance from other genotype 1a adaptive mutations in NS3 that were previously described. The contribution of the NS5A adaptive mutations to the replication of genotype 1a RNA appears to be additive to that of the NS3/4A mutations and not synergistic as shown for the combination of Q1067R and K1691R. The invention also includes methods for making replication competent polynucleotides, identifying a compound that inhibits replication of a replication competent polynucleotide, selecting a replication competent polynucleotide, and detecting a replication competent polynucleotide.

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=> D L10 IBIB BAS
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'BAS' IS NOT A VALID FORMAT FOR FILE 'CAPLUS'

The following are valid formats:

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ABS ----- GI and AB
ALL ----- BIB, AB, IND, RE
APPS ----- AI, PRAI
BIB ----- AN, plus Bibliographic Data and PI table (default)
CAN ----- List of CA abstract numbers without answer numbers
CBIB ----- AN, plus Compressed Bibliographic Data
CLASS ----- IPC, NCL, ECLA, FTERM
DALL ----- ALL, delimited (end of each field identified)
DMAX ----- MAX, delimited for post-processing
FAM ----- AN, PI and PRAI in table, plus Patent Family data
FBIB ----- AN, BIB, plus Patent FAM
IND ----- Indexing data
IPC ----- International Patent Classifications
MAX ----- ALL, plus Patent FAM, RE
PATS ----- PI, SO
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SAM ----- CC, SX, TI, ST, IT SCAN ----- CC, SX, TI, ST, IT (random display, no answer numbers; SCAN must be entered on the same line as the DISPLAY, e.g., D SCAN or DISPLAY SCAN) STD ----- BIB, CLASS IABS ----- ABS, indented with text labels IALL ----- ALL, indented with text labels IBIB ----- BIB, indented with text labels IMAX ----- MAX, indented with text labels ISTD ----- STD, indented with text labels OBIB ----- AN, plus Bibliographic Data (original) OIBIB ----- OBIB, indented with text labels SBIB ----- BIB, no citations SIBIB ----- IBIB, no citations HIT ----- Fields containing hit terms HITIND ----- IC, ICA, ICI, NCL, CC and index field (ST and IT)

containing hit terms

HITRN ----- HIT RN and its text modification

HITSTR ----- HIT RN, its text modification, its CA index name, and

its structure diagram

HITSEQ ----- HIT RN, its text modification, its CA index name, its

structure diagram, plus NTE and SEQ fields

FHITSTR ---- First HIT RN, its text modification, its CA index name, and

its structure diagram

FHITSEQ ---- First HIT RN, its text modification, its CA index name, its

structure diagram, plus NTE and SEQ fields

KWIC ----- Hit term plus 20 words on either side

OCC ----- Number of occurrence of hit term and field in which it occurs

To display a particular field or fields, enter the display field codes. For a list of the display field codes, enter HELP DFIELDS at an arrow prompt (=>). Examples of formats include: TI; TI, AU; BIB, ST; TI, IND; TI, SO. You may specify the format fields in any order and the information will be displayed in the same order as the format specification.

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L10 ANSWER 1 OF 1 CAPLUS COPYRIGHT 2008 ACS on STN

2005:523226 CAPLUS ACCESSION NUMBER:

143:54458 DOCUMENT NUMBER:

TITLE: Replication competent hepatitis C virus genotype 1a

with adaptive mutations and methods of use for drug

screening and selection of host cell line

INVENTOR(S): Lemon, Stanley M.; Yi, Minkyung

PATENT ASSIGNEE(S): Board of Regents, the University of Texas System, USA

SOURCE: PCT Int. Appl., 102 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

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L11 ANSWER 1 OF 1 CAPLUS COPYRIGHT 2008 ACS on STN

ACCESSION NUMBER: 2005:523226 CAPLUS

143:54458 DOCUMENT NUMBER:

TITLE: Replication competent hepatitis C virus genotype 1a with adaptive mutations and methods of use for drug

screening and selection of host cell line

Lemon, Stanley M.; Yi, Minkyung INVENTOR(S):

Board of Regents, the University of Texas System, USA PATENT ASSIGNEE(S):

SOURCE: PCT Int. Appl., 102 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

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a coding sequence encoding a hepatitis C virus polyprotein having adaptive mutations. The genotype 1a adaptive mutations identified here can be grouped functionally into two groups: K2040R, F2080V, and S2204I, which are all located within NS5A, and Q1067R, G1188R, V1655I, and K1691R (in NS4A), which are all located in or associated with the protease domain of NS3. These NS3 and NS4A mutations are located at some distance from other genotype 1a adaptive mutations in NS3 that were previously described. The contribution of the NS5A adaptive mutations to the replication of genotype 1a RNA appears to be additive to that of the NS3/4A mutations and not synergistic as shown for the combination of Q1067R and K1691R. The invention also includes methods for making replication competent polynucleotides, identifying a compound that inhibits replication of a replication competent polynucleotide, and detecting a replication competent polynucleotide, and detecting a replication competent polynucleotide.

=> D L12 IBIB ABS

L12 ANSWER 1 OF 1 CAPLUS COPYRIGHT 2008 ACS on STN

ACCESSION NUMBER: 2005:523226 CAPLUS

DOCUMENT NUMBER: 143:54458

TITLE: Replication competent hepatitis C virus genotype 1a

with adaptive mutations and methods of use for drug

screening and selection of host cell line

INVENTOR(S): Lemon, Stanley M.; Yi, Minkyung

PATENT ASSIGNEE(S): Board of Regents, the University of Texas System, USA

SOURCE: PCT Int. Appl., 102 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

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		AZ,	BY,	KG,	KΖ,	MD,	RU,	ΤJ,	TM,	AT,	BE,	BG,	CH,	CY,	CZ,	DE,	DK,
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		RO,	SE,	SI,	SK,	TR,	BF,	ВJ,	CF,	CG,	CI,	CM,	GA,	GN,	GQ,	GW,	ML,
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AB The invention provides replication competent polynucleotides that include a coding sequence encoding a hepatitis C virus polyprotein having adaptive mutations. The genotype 1a adaptive mutations identified here can be grouped functionally into two groups: K2040R, F2080V, and S2204I, which are all located within NS5A, and Q1067R, G1188R, V1655I, and K1691R (in NS4A), which are all located in or associated with the protease domain of NS3. These NS3 and NS4A mutations are located at some distance from other

genotype 1a adaptive mutations in NS3 that were previously described. The contribution of the NS5A adaptive mutations to the replication of genotype 1a RNA appears to be additive to that of the NS3/4A mutations and not synergistic as shown for the combination of Q1067R and K1691R. The invention also includes methods for making replication competent polynucleotides, identifying a compound that inhibits replication of a replication competent polynucleotide, and detecting a replication competent polynucleotide.

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L13 ANSWER 1 OF 1 CAPLUS COPYRIGHT 2008 ACS on STN

ACCESSION NUMBER: 2005:523226 CAPLUS

DOCUMENT NUMBER: 143:54458

TITLE: Replication competent hepatitis C virus genotype 1a

with adaptive mutations and methods of use for drug

screening and selection of host cell line

INVENTOR(S): Lemon, Stanley M.; Yi, Minkyung

PATENT ASSIGNEE(S): Board of Regents, the University of Texas System, USA

SOURCE: PCT Int. Appl., 102 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent LANGUAGE: English

FAMILY ACC. NUM. COUNT: 1

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AB The invention provides replication competent polynucleotides that include a coding sequence encoding a hepatitis C virus polyprotein having adaptive mutations. The genotype la adaptive mutations identified here can be grouped functionally into two groups: K2040R, F2080V, and S2204I, which are all located within NS5A, and Q1067R, G1188R, V1655I, and K1691R (in NS4A), which are all located in or associated with the protease domain of NS3. These NS3 and NS4A mutations are located at some distance from other genotype la adaptive mutations in NS3 that were previously described. The contribution of the NS5A adaptive mutations to the replication of genotype la RNA appears to be additive to that of the NS3/4A mutations and not synergistic as shown for the combination of Q1067R and K1691R. The invention also includes methods for making replication competent polynucleotides, identifying a compound that inhibits replication of a

replication competent polynucleotide, selecting a replication competent polynucleotide, and detecting a replication competent polynucleotide.